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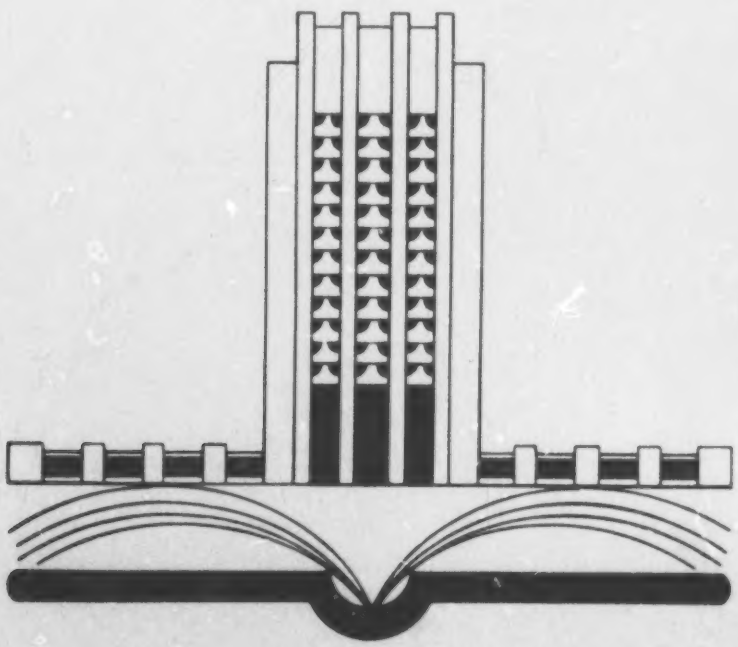

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SEARCH STRATEGY

Line	Command
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2.	S RFLP? OR RESTRICTION()FRAGMENT()LENGTH()POLYMORPHISM?
3.	S QTL? OR QUANTITATIVE()TRAIT()LOC?
4.	S GENETIC()LINKAGE?
5.	S S1 OR S2 OR S3 OR S4
6.	S S5 AND SH=L200

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Isolation and characterization of a region of the Drosophila genome which contains a cluster of differentially expressed maternal genes (yema gene region).
DEBIA. Ait-Ahmed, O. ; Thomas-Cavallin, M.-Rosset, R. Duluth, Minn. : Academic Press. Developmental biology. July 1987. v. 122 (1). p. 153-162. ill. Includes references. (NAL Call No.: DNAL 442.8 D49).

0069
Isolation of a structural gene mapping to subregions 63F of Drosophila melanogaster and 90B of Drosophila hydei polytene chromosomes.
Izquierdo, M. ; Arribas, C.~Alonso, C. Berlin, Springer. Chromosoma. 1981. v. 83 (3). p. 353-366. ill. 36 ref. (NAL Call No.: 442.8 C46).

0070
Isolation of Drosophila mutants with abnormal proboscis extension reflex.
JEZOA. Kimura, K.I. ; Shimozawa, T.~Tanimura, T. New York, N.Y. : Alan R. Liss. Journal of experimental zoology. Sept 1986. v. 239 (3). p. 393-399. ill. Includes references. (NAL Call No.: DNAL 410 J825).

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Joint segregation of biochemical loci in salmonidae. III. Linkage associations in Salmonidae including data from rainbow trout (Salmo gairdneri) (Comparative gene mapping).
May, B. ; Wright, J.E. Jr.~Johnson, K.R. New York, Plenum Press. Biochemical genetics. Feb 1982. v. 20 (1/2). p. 29-40. Includes 2 p. ref. (NAL Call No.: QR73.B5).

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Linkage groups of the horse (Equus caballus).
Weitkamp, L.R. ; Anderson, L. Cold Spring Harbor, N.Y. : Cold Spring Harbor Laboratory. Genetic maps. 1984. v. 3. p. 390-391. Includes 17 references. (NAL Call No.: DNAL QH445.2.G4).

0073
Linkage map of the yellow fever mosquito Aedes aegypti (2 N=6).
Munstermann, L.E. Cold Spring Harbor, N.Y. : Cold Spring Harbor Laboratory. Genetic maps. 1984. v. 3. p. 324-329. Includes 71 references. (NAL Call No.: DNAL QH445.2.G4).

0074
Linkage of albino and hemoglobin beta-chain loci in the rabbit.
JOHEA. Sandberg, K. ; Andersson, L. Washington, D.C. : American Genetic Association. The Journal of heredity. Mar/Apr 1987. v. 78 (2). p. 124-125. Includes references. (NAL Call No.: DNAL 442.8 AM3).

0075
Linkage studies in Tribolium castaneum (Herbst). XII. A revision of linkage group II.
GENDE3. Sokoloff, A. ; Ferrone, R.F.~Chaney, J.D.~Braden, J.~Munoz, R.J. Ottawa : National Research Council of Canada. Genome. Feb 1987. v. 29 (1). p. 26-33. Includes references. (NAL Call No.: DNAL QH431.G452).

0076
Linked loci of the chicken-Gallus gallus (G. domesticus).
Somes, R.G. Cold Spring Harbor, N.Y. : Cold Spring Harbor Laboratory. Genetic maps. 1984. v. 3. p. 465-473. Includes 123 references. (NAL Call No.: DNAL QH445.2.G4).

0077
Mapping four genes coding for biochemical characters on the second chromosome of Drosophila imeretensis.
DKBSAS. Goncharenko, G.G. ; Mitrofanov, V.G.~Korochkin, L.I. New York, N.Y. : Consultant Bureau. Doklady : biological sciences - Akademiia nauk SSSR. Translated from: Akademiia nauk SSSR Doklady. (3) p. 732-735. (511 P444A). May/June 1984. v. 276 (1/6). p. 349-352. ill. Includes 10 references. (NAL Call No.: DNAL 511 P444AEB).

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Mapping genes in domesticated animals.
Ruddle, F.H. ; Fries, R. New York : Plenum Press, c1986. Genetic engineering of animals : an agricultural perspective / edited by J. Warren Evans and Alexander Hollaender ; technical editor, Claire M. Wilson. Paper presented at a symposium on "Genetic Engineering of Animals," September 9-12, 1985, Davis, California. p. 39-57. ill. Includes 61 references. (NAL Call No.: DNAL SF756.5.S97 1985).

0079
Mapping of C2, Bf, and C4 genes to the swine major histocompatibility complex (swine leukocyte antigen).
JOIMA3. Lie, W.R. ; Rothschild, M.F.~Warner, C.M. Baltimore, Md. : The American Association of Immunologists. The Journal of immunology. Nov 1987. v. 139 (10). p. 3388-3395. ill. Includes references. (NAL Call No.: DNAL 448.8 J8232).

0080
Mapping of the gene for G blood group antigens to chromosome number 15 in swine.
Maurer, R.R. ; Fries, H.R. Beltsville, Md. : The Service, ARS - United States Department of Agriculture, Agricultural Research Service. In subseries: Swine Research. Progress Report No. 2. Nov 1986. (54). p. 39-40. ill. (NAL Call No.: DNAL aS21.R44A7).

0081
Mapping of the genome of the silver fox. I. Determination of chromosome localization of eight genes of the fox and a search for homologous regions of fox and human chromosomes.
SOGEBZ. Rubtsov, N.B. ; Grafodatskii, A.S.~Matveeva, V.G.~Nesterova, T.B.~Kul'bakina, N.A.~Zakiyan, S.M. New York, N.Y. : Consultants Bureau. Soviet genetics. Translated from Genetika, v. 24 (1), 1988, p. 69-78. (QH431.A1G4). July 1988. v. 24 (1). p. 44-51. ill. Includes references. (NAL Call No.: DNAL QH431.A1G43).

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A method for mapping intranuclear protein-DNA interactions and its application to a nuclease hypersensitive site.
PNASA. Jackson, P.D. ; Felsenfeld, G. Washington, D.C. : The Academy. Proceedings of the National Academy of

Sciences of the United States of America. Apr 1985. v. 82 (8). p. 2296-2300. ill. Includes 24 references. (NAL Call No.: DNAL 500 N21P).

0083
Micronuclear DNA sequences from Tetrahymena do not confer mitotic stability on ARS plasmids in Saccharomyces.
GENOE3. Horsfall, W.H. ; Pearlman, R.E. Ottawa : National Research Council of Canada. Genome. Oct 1988. v. 30 (5). p. 690-696. Includes references. (NAL Call No.: DNAL QH431.G452).

0084
Molecular analysis of the P-M gonadal dysgenesis cline in Eastern Australian Drosophila melanogaster.
GENTA. Boussy, I.A. ; Healy, M.J.~Oakeshott, J.G.~Kidwell, M.G. Baltimore, Md. : Genetics Society of America. Genetics. Aug 1988. v. 119 (4). p. 889-902. ill. Includes references. (NAL Call No.: DNAL 442.8 G28).
The latitudinal cline in P-M gonadal dysgenesis potential in easternAustralia has been shown to comprise three regions which are, from north to south respectively, P,Q, and M, with the P-to-Q and Q-to-M transitions occurring over relatively short distances. The P element complements of 30 lines from different regions of the cline were determined by molecular techniques. The total amount of P element-hybridizing DNA was high in all lines, and it did not correlate in any obvious way with the P-M phenotypes of individual lines. The number of potentially full-sized P elements per genome was high in lines from the P regions, but variable or low among lines from the Q and M regions, and thus declined overall from north to south. A particular P element deletion-derivative, the KP element, occurred in all the tested lines. The number of KP elements was low in lines from the P region, much higher in lines from the Q region, and highest among lines from the M region, thus forming a cline reciprocal to that of the full-sized P elements. Another transposable element, hobo, which has been described as causing dysgenic traits similar to those of P-M hybrid dysgenesis, was shown to be present in all lines and to vary among them in number, but not in any latitudinal pattern. The P-M cline in gonadal dysgenesis potential can be inferred to be based on underlying clinal patterns of genomic P element complements. P activity of a line was positively correlated with the number of full-sized P elements in the line, and negatively correlated with the number of KP elements. Among Q and M lines, regulatory ability was not correlated with numbersof KP elements.

0085
Molecular and genetic characterization of the Drosophila melanogaster 87E actin gene region.
GENTA. Manseau, L.J. ; Ganetzky, B.~Craig, E.A. Baltimore, Md. : Genetics Society of America. Genetics. June 1988. v. 119 (2). p. 407-420. ill. Includes references. (NAL Call No.: DNAL 442.8 G28).
Abstract: A combined molecular and genetic analysis of the 87E actin gene (Act87E) in Drosophila melanogaster was undertaken. A clone of Act87E was isolated and characterized. The Act87E transcription unit is 1.57 kb and includesa 556-base intervening sequence in the 5' leader of the gene. The protein-codingregion is contiguous and encodes a protein that is >93% identical to the other Drosophila actins. By in situ hybridization with a series of deficiencies that break in 87E, Act87E was localized to a region encompassing one to three faint, polytene chromosome bands. The region between the deficiency endpoints that flank the actin gene was isolated and measures approximately 24-30 kb. The closest proximal deficiency endpoint lies8-10 kb 5' to the actin gene; the closest distal deficiency endpoint lies 16-20 kb 3' to the actin gene. A single, recessive lethal complementation group lies between the deficiency endpoints that flank the actin gene. An EMS mutagenesis screen produced four additional members of this recessive lethal complementation group. Molecular analysis of the members of this complementation group indicated that two of the newly induced mutations have deletions of approximately 1 kb in a transcribed region 4-5 kb 3' (distal) to the actin gene. This result suggests that the recessive lethal complementation group represents a gene separate from and distal to the actin gene. The mutagenesis screen failed to identify additional recessive lethal complementation groups in the actin gene-containing region. The implications of the failure to identify recessive lethal mutations in the actin gene are discussed in reference to studies of other conserved mutigene families and other muscle protein mutations.

0086
Molecular cytogenetics of cattle: a genomic approach to disease resistance and productivity.
JDSCAE. Womack, J.E. Champaign, Ill. : American Dairy Science Association. Journal of dairy science. Apr 1988. v. 71 (4). p. 1116-1123. ill. Includes references. (NAL Call No.: DNAL 44.8 J822).

0087
Molecular markers in the genetic improvement of farm animals.
Beckmann, J.S. ; Soller, M. New York, N.Y. : Nature Pub. Co. Bio/technology. June 1987. v. 5 (6). p. 573-576. ill. Includes references. (NAL Call No.: DNAL QH442.B5).

0088
Mutations affecting expression of the rosy locus in Drosophila melanogaster.
GENTA. Lee, C.S. ; Curtis, D.~McCarron, M.~Love, C.~Gray, M.~Bender, W.~Chovnick, A. Baltimore, Md. : Genetics Society of America. Genetics.

May 1987. v. 116 (1). p. 55-66. ill. Includes references. (NAL Call No.: DNAL 442.8 G28).

0089
A new era in mammalian gene mapping: somatic cell genetics and recombinant DNA methodologies.
Ruddle, F.H. London, Macmillan Journals. Nature. Nov 12/18, 1981. v. 294 (5837). p. 115-120. Includes 67 ref. (NAL Call No.: 472 N21).

0090
Organization and expression of a second chromosome follicle cell gene cluster in Drosophila.
DEBIA. Popodi, E. ; Minoo, P.~Burke, T.~Waring, G.L. Duluth, Minn. : Academic Press. Developmental biology. June 1988. v. 127 (2). p. 248-256. Includes references. (NAL Call No.: DNAL 442.8 D49).

0091
Partial molecular genetic map of the rabbit VH chromosomal region.
JOIMA3. Currier, S.J. ; Gallarda, J.L.~Knight, K.L. Baltimore, Md. : The American Association of Immunologists. The Journal of immunology. Mar 1, 1988. v. 140 (5). p. 1651-1659. ill. Includes references. (NAL Call No.: DNAL 448.8 J8232).

0092
Provisional assignment of the G blood-group locus to chromosome 15 in swine--gene mapping in swine using natural and induced marker chromosomes.
Fries, R.JOHEA. ; Stranzinger, G.~Vogeli, P. Washington, D.C. : American Genetic Association. The Journal of heredity. Nov/Dec 1983. v. 74 (6). p. 426-430. Includes references. (NAL Call No.: 442.8 AM3).

0093
Rates of nuclear DNA evolution in pheasant-like birds: evidence from restriction maps.
PNASA. Helm-Bychowski, K.M. ; Wilson, A.C. Washington, D.C. : The Academy. Proceedings of the National Academy of Sciences of the United States of America. Feb 1986. v. 83 (3). p. 688-692. ill. Includes references. (NAL Call No.: DNAL 500 N21P).

0094
A rehabilitation of the genetic map of the 84B-D region in drosophila melanogaster.
GENTA. Cavener, D.R. ; Otteson, D.C.~Kaufman, T.C. Baltimore, Md. : Genetics Society of America. Genetics. Sept 1986. v. 114 (1). p. 111-123. ill. Includes references. (NAL Call No.: DNAL 442.8 G28).

0095
Relative positions of the loci of the peacomb (P), eumelanin restrictor (Db), eumelanin extension (Ml) and plumage pattern (Pg) genes of the domestic fowl.
BPOSA. Carefoot, W.C. Oxfordshire : Carfax Publishing Company. British poultry science. June 1987. v. 28 (2). p. 347-350. Includes references. (NAL Call No.: DNAL 47.8 B77).

0096
Restricted ovulators linked to heart disease and gene mapping (Laying hens).
Laughren, C.M.POTRA. Mount Morris : Watt Publishing Co. Poultry tribune. Nov 1983. v. 89 (11). p. 37-38. ill. (NAL Call No.: 47.8 P8642).

0097
Restriction fragment length polymorphism among Israeli Holstein-Friesian dairy bulls.
Beckmann, J.S. ; Kashi, Y.~Hallerman, E.M.~Nave, A.~Soller, M. Oxford : Blackwell Scientific Publications. Animal genetics. 1986. v. 17 (1). p. 25-38. ill. Includes references. (NAL Call No.: DNAL QP98.A1A5).

0098
Restriction fragment length polymorphism (RFLP) analysis of bovine nuclear protein genes.
THAGA. Krawetz, S.A. ; Bricker, R.A.~Connor, W.~Church, R.B.~Dixon, G.H. Berlin, W. Ger. : Springer International. Theoretical and applied genetics. 1988. v. 75 (3). p. 402-409. ill. Includes references. (NAL Call No.: DNAL 442.8 Z8).

0099
Restriction fragment length polymorphisms in poultry breeding.
POSCAL. Soller, M. ; Beckmann, J.S. Champaign, Ill. : Poultry Science Association. Poultry science. Aug 1986. v. 65 (8). p. 1474-1488. Includes references. (NAL Call No.: DNAL 47.8 AM33P).

0100
Restriction fragment length polymorphisms in varietal identification and genetic improvement /Jacques S. Beckmann, Morris Soller, Benjamin Burr.
Beckmann, Jacques S. ; Soller, Morris.~Burr, Benjamin. Bet Dagan, Israel : BARD, 1986. Final report.~"Project no. I-397-82R.". 118 p. : ill. ; 28 cm. Includes bibliographies. (NAL Call No.: DNAL QH442.B4).

0101
The rosy region of Drosophila melanogaster and Drosophila simulans. I. Contrasting levels of naturally occurring DNA restriction map variation and divergence.
GENTA. Aquadro, C.F. ; Lado, K.M.~Noon, W.A. Baltimore, Md. : Genetics Society of America. Genetics. Aug 1988. v. 119 (4). p. 875-888. Includes references. (NAL Call No.: DNAL 442.8 G28).
A 40-kb region around the rosy and snake loci was analyyzed for restriction map variation among 60 lines of Drosophila melanogaster and 30 lines of Drosophila simulans collected together at a single locality in Raleigh, North Carolina. DNA sequence variation in D. simulans was estimated to be 6.3 times greater than in D. melanogaster (heterozgosities per nucleotide of 1.9% vs. 0.3%). This result stands in marked contrast to

results of studies of phenotypic variation including proteins (allozymes), morphology and chromosome arrangements which are generally less variable and less geographically differentiated in D. simulans. Intraspecific polymorphism is not distributed uniformly over the 40-kb region. The level of heterozygosity per nucleotide varies more than 12-fold across the region in D. simulans, being highest over the hsc2 gene. Similar, though less extreme, variation in heterozygosity is also observed in D. melanogaster. Average interspecific divergence (corrected for intraspecific polymorphism) averaged 3.8%. The pattern of interspecific divergence over the 40-kb region shows some disparities with the spatial distribution of intraspecific variation, but is generally consistent with selective neutrality predictions: the most polymorphic regions within species are generally the most divergent between species. Sequence-length polymorphism is observed for D. melanogaster to be at levels comparable to other gene regions in this species. In contrast, no sequence length variation was observed among D. simulans chromosomes (limit of resolution approximately 100 bp). These data indicate that transposable elements play at best a minor role in the generation of naturally occurring genetic variation in D. simulans compared to D. melanogaster. We hypothesize that differences in species effective population size are the major determinant of the contrasting levels and patterns of DNA sequence and insertion/deletion variation that we report here and the patterns of allozym

0102
A selective screen to recover chromosomal deletions and duplications in Drosophila melanogaster.
GENTA. Gubb, D. ; McGill, S.~Ashburner, M. Baltimore, Md. : Genetics Society of America. Genetics. June 1988. v. 119 (2). p. 377-390. ill. Includes references. (NAL Call No.: DNAL 442.8 G28).
Abstract: A screen is described that will select for breakpoints within a restricted chromosomal region in Drosophila. The aberrations recovered can be used to construct chromosomes carrying synthetic duplications and deletions. Such chromosomes have applications in the mapping of complementation groups at both the genetic and molecular level. In particular, breakpoints recovered after P element hybrid dysgenesis tend to be associated with P element insertion sites. Such aberration breakpoints can be genetically mapped, as synthetic deletions,and then used as transposon-tagged sites for the recovery of genomic clones.

0103
Sequence of the viral replicase gene from foot-and-mouth disease virus C1-Santa Pau (C-S8).
GENED6. Martinez-Salas, E. ; Ortin, J.~Domingo, E. Amsterdam : Elsevier Science Publishers. Gene. 1985. v. 35 (1/2). p. 55-61. ill. Includes references. (NAL Call No.: DNAL QH442.A1G4).

0104
Short-antenna, a new mutant in the mosquito Aedes (Finlaya) togoi (Genetic mapping).
Tadano, T.ESDBA. Tokyo, Nihom Eisei Dobutsu Gakkai. Eisei dobutsu = Japanese journal of sanitary zoology. June 15, 1982. v. 33 (2). p. 105-109. ill. 1 p. ref. (NAL Call No.: QL99.E3).

0105
Somatic cell genetics and animal breeding.
Forster, M. Oxford : Oxford University Press, c1986. Exploiting new technologies in animal breeding : genetic developments / edited by C. Smith, J.W.B. King, and J.C. McKay. p. 47-53. Includes references. (NAL Call No.: DNAL SF105.E9).

0106
Structure of chick chromosomal genes for calbindin and calretinin.
JMOBAK. Wilson, P.W. ; Rogers, J.~Harding, M.~Pohl, V.~Pattyn, G.~Lawson, D.E.M. London : Academic Press. Journal of molecular biology. Apr 20, 1988. v. 200 (4). p. 615-625. ill. Includes references. (NAL Call No.: DNAL 442.8 J8224).

0107
Structure of the chicken link protein gene: exons correlate with the protein domains.
PNASA. Kiss, I. ; Deak, F.~Mestric, S.~Delius, H.~Soos, J.~Dekany, K.~Argraves, W.S.~Sparks, K.J.~Goetinck, P.F. Washington, D.C. : The Academy. Proceedings of the National Academy of Sciences of the United States of America. Sept 1987. v. 84 (18). p. 6399-6403. ill. Includes references. (NAL Call No.: DNAL 500 N21P).

0108
Structure of the chromosomal chicken progesterone receptor gene.
PNASA. Huckaby, C.S. ; Conneely, O.M.~Beattie, W.G.~Dobson, A.D.W.~Tsai, M.J.~O'Malley, B.W. Washington, D.C. : The Academy. Proceedings of the National Academy of Sciences of the United States of America. Dec 1987. v. 84 (23). p. 8380-8384. ill. Includes references. (NAL Call No.: DNAL 500 N21P).

0109
Structure of the cloned Locusta migratoria mitochondrial genome: restriction mapping and sequence of its ND-1 (URF-1) gene.
CUGEDS. McCracken, A. ; Uhlenbusch, I.~Gellissen, G. Berlin, W. Ger. : Springer International. Current genetics. 1987. v. 11 (8). p. 625-630.

ill. Includes references. (NAL Call No.: DNAL QH426.C8).

O110
Structure of the polymorphic feline c-myc oncogene locus.
GENED6. Soe, L.H. ; Roy-Burman, P. Amsterdam : Elsevier Science Publishers. Gene. Nov 1984. v. 31 (1/3). p. 123-128. ill. Includes references. (NAL Call No.: DNAL QH442.A1G4).

O111
Studies of the genetic organization of the vestigial microregion of Drosophila melanogaster.
GENTA. Lasko, P.F. ; Pardue, M.L. Baltimore, Md. : Genetics Society of America. Genetics. Oct 1988. v. 120 (2). p. 495-501. Includes references. (NAL Call No.: DNAL 442.8 G28).
The region of the second chromosome of Drosophila melanogaster defined by Df(2R)vgB was screened for recessive lethal and visible mutations. Fifty-eight new recessive alleles fall into 17 complementation groups. Many new vg alleles were also isolated in a screen for new vg deficiencies. The breakpoints of the new vg deficiencies were nonrandomly distributed. The distal breakpoints of twelve of 20 deficiencies overlapping Df(2R)vgB are genetically identical to that of Df(2R)vgD, coinciding with the position of a complex, pleiotropic locus, l(2)49Ea-Psc-Su(z)2.

O112
Studies on genetic linkage in domestic animals with special reference to the horse /Leif Andersson. --.
Andersson, Leif. Uppsala, Sweden : Dept. of Animal Breeding and Genetics, Swedish University of Agricultural Sciences, 1983. Thesis (doctoral)--Sveriges Lantbruks-universitet, 1983. Spine title: Genetic linkage in domestic animals. 1 v. (various pagings) : col. ill. ; 25 cm. Includes bibliographies. (NAL Call No.: DNAL QH432.A4).

O113
Suppressible P-element alleles of the vestigial locus in Drosophila melanogaster.
MGGEAE. Williams, J.A. ; Pappu, S.S.~Bell, J.B. Berlin, W. Ger. : Springer International. M G G : Molecular and general genetics. May 1988. v. 212 (2). p. 370-374. ill. Includes references. (NAL Call No.: DNAL 442.8 Z34).

O114
Tissue-specific and constitutive alpha-tubulin genes of Drosophila melanogaster code for structurally distinct proteins.
PNASA. Theurkauf, W.E. ; Baum, H.~Bo, J.~Wensink, P.C. Washington, D.C. : The Academy. Proceedings of the National Academy of Sciences of the United States of America. Nov 1986. v. 83 (22). p. 8477-8481. ill. Includes references. (NAL Call No.: DNAL 500 N21P).

O115
Toward a physical map of the genome of the nematode Caenorhabditis elegans.
PNASA. Coulson, A. ; Sulston, J.~Brenner, S.~Karn, J. Washington, D.C. : The Academy. Proceedings of the National Academy of Sciences of the United States of America. Oct 1986. v. 83 (20). p. 7821-7825. ill. Includes references. (NAL Call No.: DNAL 500 N21P).

O116
Use of peptide tagging to detect proteins expressed from cloned genes: deletion mapping functional domains of Drosophila hsp70.
EMJODG. Munro, S. ; Pelham, H.R.B. Oxford, Eng. : IRL Press. The EMBO journal - European Molecular Biology Organization. Dec 20, 1984. v. 3 (13). p. 3087-3093. ill. Includes references. (NAL Call No.: DNAL QH506.E46).

O117
Y chromosome loops in Drosophila melanogaster.
GENTA. Bonaccorsi, S. ; Pisano, C.~Puoti, F.~Gatti, M. Baltimore, Md. : Genetics Society of America. Genetics. Dec 1988. v. 120 (4). p. 1015-1034. ill. Includes references. (NAL Call No.: DNAL 442.8 G28).

O118
7C female sterile mutants fail to accumulate early eggshell proteins necessary for later chorion morphogenesis in Drosophila.
DEBIA. Bauer, B.J. ; Waring, G.L. Duluth, Minn. : Academic Press. Developmental biology. June 1987. v. 121 (2). p. 349-358. Includes references. (NAL Call No.: DNAL 442.8 D49).

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